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		Il ial		BoCAL Priner	13 100.0t	RCORV	٠.				•	
	1	CTAGATAACG	AGGGCAAAAA	ATGAAAAGA	CAGCTATCGC	GATTGCAGTG	GCACTGGCTG	GITTOGCTAC	CGTAGGGCAG	GCCGATATCG	TGCTGACCCA	
•	101	GAGCCCCGGCG	ACCCTGAGCC	TACTITITICI TGTCTCCGGG	CGAACGTGCG	ACCCTGAGCT	GCAGAGCGAG	CCAGTCTGTT	TCTCGTTCTT	ATCTGGCTTG	GTACCAGCAG	
	201	CTCGGGCCGC AAACCAGGTC	TGGGACTCGG AAGCACCGCG	ACAGAGGCCC TCTATTAATT	GCTTGCACGC TATGGTGCTT	TGGGACTCGA	CGTCTCGCTC AACTGGGGTC	GGTCAGACAA CCGGCGCGTT	AGAGCAAGAA TTAGCCCCCCCC	TAGACCGAAC	CATGGTCGTC	
		TTTGGTCCAG	TTCGTGGCGC Bbs1	AGATAATTAA	ATACCACGAA MscI	GAGCAGCACG	TTGACCCCAG	CCCCCCCCAA	AATCGCCGAG	ACCTAGGCCG	TGCCTAAAAT	
	201	000003.003.00	***	01.3.00man sia		mma mma maga	01 001 00000					
	301	OGGACTOGTA	ATCGTCGGAC	CTTGGACTTC	TGAAACGCTG	AATAATAACG	CAGCAGCGTG GTCGTCGCAC	CATTAATAAG	TATTACCTTT ATAATGGAAA	GECCAGGETA CCGGTCCCAT	CGAAAGTTGA GCTTTCAACT	
•		BSINI		•	· · · · · · · · · · · · · · · · · · ·	• .	• . • .		٠		•	
	401	AATTAAACGT	ACCGTCGCTC	CTCCGAGCGT GAGGCTCGCA	GTTATTITT	COCCGAGOG	ATGAACAACT	GAAAAGCGGC	ACGCCCACCC	TOGTGTGCCT	GCTGAACAAC	
	501	TTTTATCCGC	GTGAAGCGAA	AGTTCAGTGG	AAAGTAGACA	ACGCCCTGCA	AAGOGGCAAC	AGCCAGGAAA	GOGTGACOGA	ACAGGATAGC	AAAGATAGCA	
	601	CCTATTCTCT	GAGCAGCACC	TCAAGTCACC CTGACCCTGA	GCAAAGCGGA	TTATGAAAAA	CATAAAGTGT	ATGCGTGCGA	AGTGACCCAT	CAAGGTCTGA	GCAGCCCCCT	
		GGATAAGAGA	CTCGTCGTCG	GACTGGGACT	CGTTTCGCCT Bucal f	AATACTITT or 100.0%	GTATTTCACA	TACGCACGCT	TCACTGGGTA	GTTCCAGACT	CGTCGGGCCA	•
	. ,	St	ul Sphi	·		SapI						
٠.				. .		*******	·	•				
	- 701	(1) (III) 1 1 III (III)		0001000000	· bol	Primer 1 100.0	}	1011100100	100001000	61 mmm 66		
	101	CTGATTTAGA	AAATTAGCAC	GCGAGGCCTG CGCTCCGGAC	TATTCGTACG	CATCCTCTT	TATTTTACTT	TGTTTCGTGA	TAACGTGACC	CACTUTTACC	CAACGAGAAG	
		ECAL for 100	.OR Mel								•	
	•				•	·			•	• • • • • •		
	801	ACCCCTGTTA	CCAAAGCCCA	GGTGCAATTG CCACGTTAAC	GTGGAAAGCG	GCGGCGGCCT	COTOCAACCG	GGCGCAGCC	TGCGTCTGAG	CTGGGGGGCC	TCCCGATTTA	
	901	CCTITTCTTC	TTATGGTGGT	AATTGGGTGC	GCCAAGCCCC	TGGGAAGGGT	CTCGAGTOGG	TGAGOGGTAT	CCATTATTCT	GGTAGCTCTA	CCTATTATGC	
	1001	GGATAGCGTG	AAAGGCCGTT	TTAACCCACG TTACCATTTC	ACGTGATAAT	TOGAAAAACA	CCCTGTATCT	GCAAATGAAC	AGCCTGCGTG	CGGAAGATAC	GGCGTGTAT	
		CCTATCGCAC	TTTCCGGCAA	AATGGTAAAG	TGCACTATTA Sali	AGCTTTTTGT	GGGACATAGA	CGTTTACTTG	TCGGACGCAC	CCCTTCTATG	CCGCCACATA	
	·	BssHII Sapi	•	Styl	BlpI			• • •			` :	
	. 1101		्राज्यम् (मार्शनिक्यसम्बद्धमारुगः)				MM137M	100000000		1000000100	1110000001	
	٠.	ATAACGCGCG	CACGAGAAGT	TAAGTGGGCT ATTCACCCGA	CCAACCCCAA	AACTAGTAAC	CCCGGTTCCG	TGGGACCACT	GCCAATCGAG	TOGCAGCTOG	TTYYYAYTT	•
: '		CGCACAAAGG	CGACCGAGGC	AGCAGCAAAA TCGTCGTTTT	CGTCGTCGCC	GCCGTGCCGA	CGGGACCCGA	CGGACCAATT	TCTAATAAAG	GEOCTIGGIC	YENCHYPA	
٠.	1301	CTGGAACAGC	GGGGGGTGA	CCAGCGGCGT GGTCGCCGCA	GCATACCTTT	CCGCCGTGC	TGCAAAGCAG	CCCCTGTAT	AGCCTGAGCA	CCGTTGTGAC	CGTCCCCGACC	
			•		Kooli	0000000		· · ·	reduncted	COCACACTO	CONSISTING	
		1000000000					· · ·	· · · ·				
	1901	DTAADOTEOT TOTAADOTEOT LIE228	GCACTCAGAC CGTGAGTCTG	CTATATTTGC GATATAAACG	AACGTGAACC TTGCACTTGG	ATAAACCGAG TATTTGGCTC	CAACACCAAA GTTGTGGTTT	GIUGATAAAA CACCIATITT	AAGRIGAACC TTCACCTRGG	GAAAAGCGAA CTTTTCGCTT	TTCCCAGGG AAGGGTCCCC	•
		Asci	Rin	dii		•			•	٠.	•	
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FIG. 2B-1

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